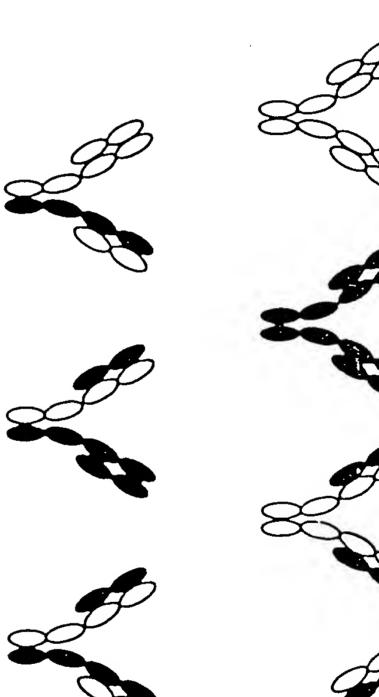
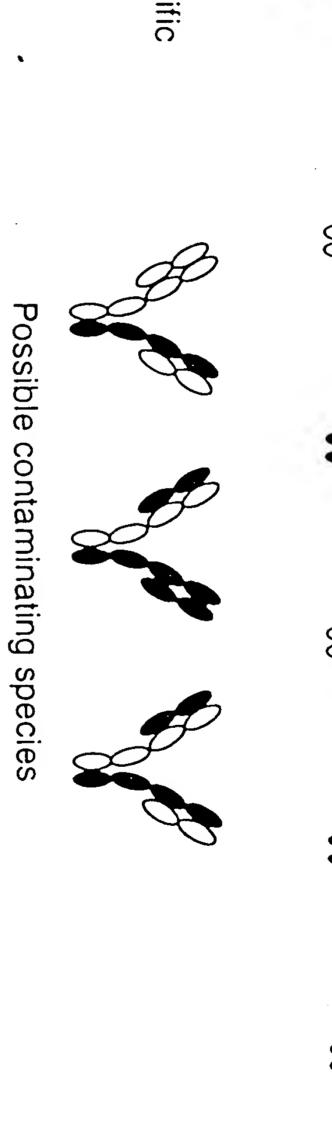
A) Before engineering of CH3 domain

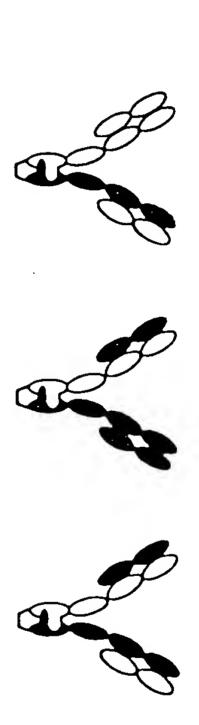


Target bispecific antibody





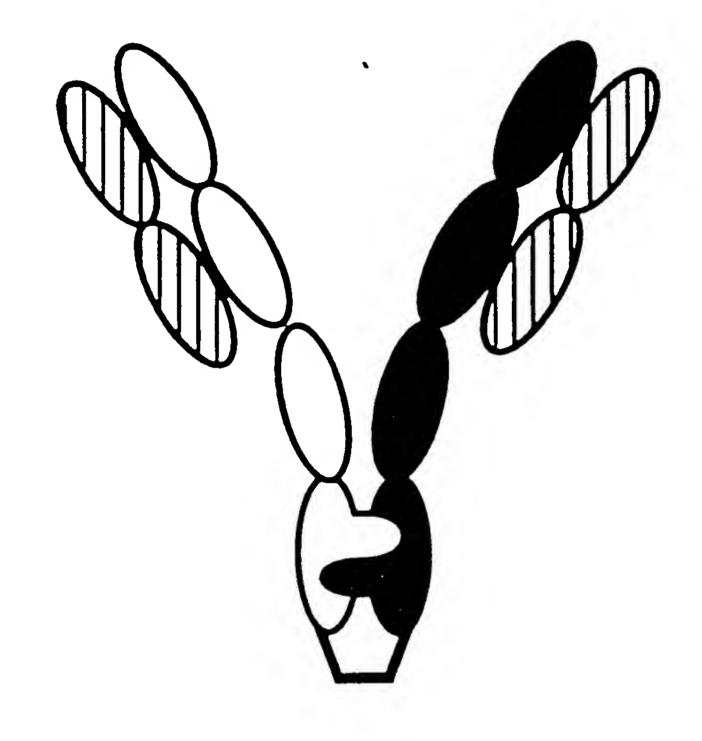
B) After engineering of CH3 domain



Possible major contaminating species

Target bispecific antibody

= Engineered disulfide bond between CH3 domains 



Target bispecific antibody

Fig. 2A

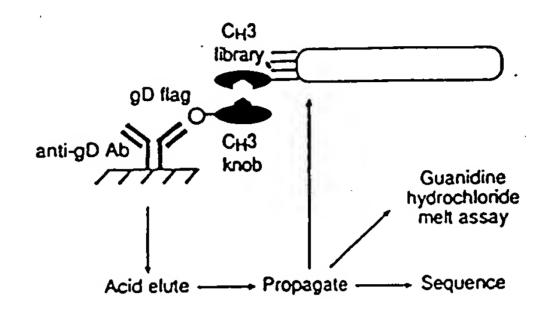


Fig. 2B

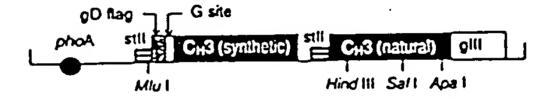
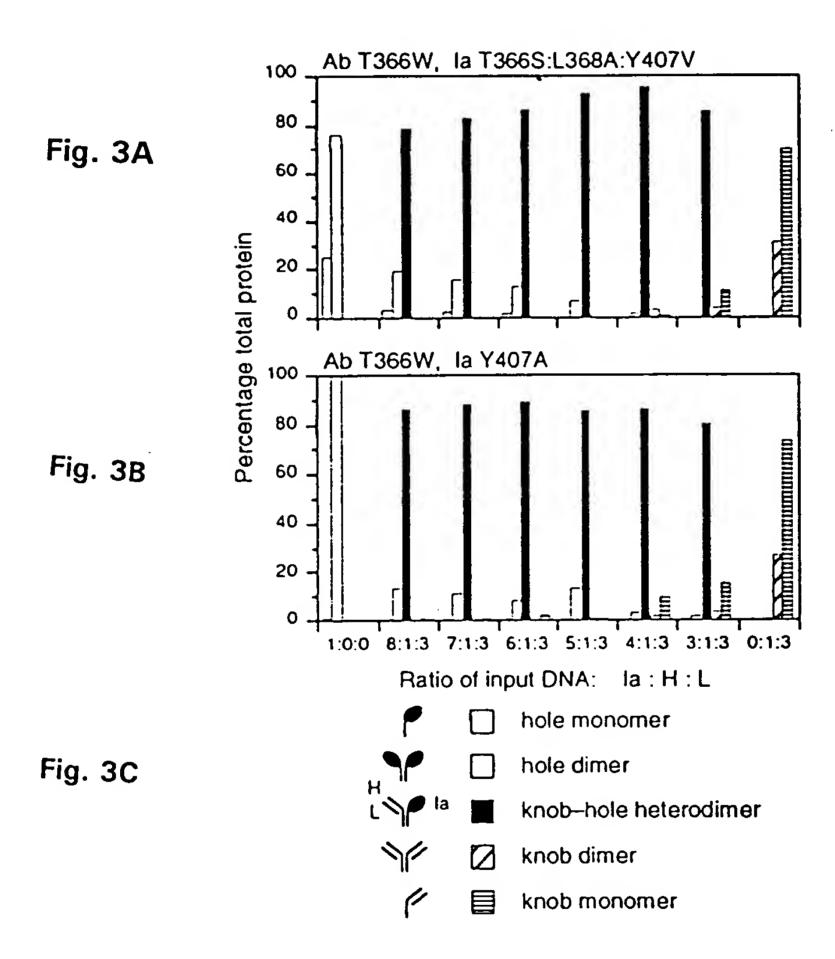


Fig. 2C

(SEQ ID NO: 13)



## 

r

50 YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV GOSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV YEGSKRPSGV	(SEQ ID NO: 14) (SEQ ID NO: 15) (SEQ ID NO: 17) (SEQ ID NO: 18) (SEQ ID NO: 19) (SEQ ID NO: 20) (SEQ ID NO: 21) (SEQ ID NO: 22) (SEQ ID NO: 22)
1 20 abc 30 40 50  QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKGPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKGPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV QSVLTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIYEGSKRPSGV CDR L1	60 90 a 100  SNRFSGSKSGNTASLTISGLQAEDEADYYCSSYTTRSTRVFGGGTKLTVL CDR L3  CDR L3
Axl.78 Rse.23 IgER.MAT2C1G11 GCSFR.A4 Rse.04 obr.4 Rse.20 Rse.15 vegf.5	Ax1.78 Rse.23 IGER.MAT2C1G11 GCSFR.A4 Rse.04 obr.4 Rse.20 Rse.15 vegf.5

$V_{H}$								
***	* *** *	* * *	30 ASGFTFS <u>SY</u> ** ** TVSGGSIS <u>SC</u> CDR	- G <u>GYYWS</u> WI	** ***	*** *	****	
* * *	*** * **	** *	abc NRLRAEDTA	*****		WYFDLWGRO	** ****	(SEQ ID NO: 23)
<u>NPSLKS</u> R	VTISVDTS	KNQFSLKI	LSSVTAADTA	AVYYCARI	VDLEDYGS	<u>GASDY</u> WGQ	GTLVTVSS	(SEQ ID NO: 24)
CDR H2					CDR	Н3		
V <sub>L</sub>								
her3.18 DIQMTQS obr.26	10 PSTLSASIO	20 GDRVTIT	30 CRASEGIYH CDR L	<u>wla</u> wyqqi	40 KPGKAPKL	5 <u>0</u> LIY <u>KASSL</u> CDR L		
	O CDFTLTISS	80 LQPDDFA	90 <u> </u>		GTKLEIK	(SEQ I	ID NO: 25)	

Percentage Identity of anti-ObR and anti-HER3  $V_{\scriptscriptstyle L}$ 

	Н1	Н2	Н3	H4	Н5	Н6	н7	Н8	н9	н10	н11
01	49	47	51	81	60	48	76	51	100	62	51
02	84	79	88	50	48	99	48	88	48	45	88
03	83	82	85	51	50	95	49	85	49	46	85
04	47	50	51	83	77	48	65	51	73	64	51
05	49	47	51	81	60	48	76	51	100	62	51
06	83	79_	86	50	50	99	47	86	48	45	86
07	81	100	86	51	49	80	48	86	47	44	86
08	81	100	86	51	49	80	48	86	47	44	86
09	81	100	86	51	49	80	48	86	47	44	86
010	83	79	85	50	49	98	46	85	48	45	85
011	83	80	87 .	50	49	99	47	87	48	45	87
012	81	100	86	51	49	80	48	86	47	44	86
013	49	47	51	81	60	48	76	51	100	62	51
014	50	50	54	95	67	49	76	54	75	62	54
015	82	79	85	49	48	97	46	85	47	44	85
016	84	80	87	50	49	100	47	87	48	45	87
017	45	44	47	65	62	45	62	47	62	100	47
018	50	51	50	75	79	50	63	50	66	62	50

01-018: Anti-Ob-R antibody clones obr. 1, 11, 12, 14, 15, 16, 17, 18, 19, 2, 20, 21, 22, 23, 24, 26, 3, 4, respectively.

H1-H11: Anti-HER3 antibody clones her3.1, 3.10, 3.11, 3.12, 3.16, 3.18, 3.19, 3.22, 3.3, 3.4, 3.7, respectively.